## **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/569,224
Source:	TEWP.
Date Processed by STIC:	03/02/2006

## ENTERED



IFWP

**RAW SEQUENCE LISTING**PATENT APPLICATION: **US/10/569,224**DATE: 03/02/2006

TIME: 10:25:02

Input Set : A:\1300-1-013PCTUS - SEQUENCE LIST.txt

Output Set: N:\CRF4\03022006\J569224.raw

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3 <110> APPLICANT: STAMPS, Alasdair
     5 <120> TITLE OF INVENTION: A PROTEIN INVOLVED IN CARCINOMA
     7 <130> FILE REFERENCE: 1300-1-013PCT/US
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/569,224
C--> 10 <141> CURRENT FILING DATE: 2006-02-22
    12 <150> PRIOR APPLICATION NUMBER: GB0320878.2
    13 <151> PRIOR FILING DATE: 2003-09-05
    15 <160> NUMBER OF SEQ ID NOS: 6
    17 <170> SOFTWARE: PatentIn version 3.1
    19 <210> SEQ ID NO! 1
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     46 Met Arg Gly Gly Ile Leu Tyr Thr Glu Pro Arg Asp Ser Met Gln Tyr
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                                    120
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    62 Val Ile Leu Ser Val Cys Ile Thr Val Leu Cys Val Phe Asp Pro Thr
     63 145
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     66 Gly Arg Thr Phe Val Lys Leu Arg Ala Thr Lys Arg Arg Gln Arg Asn
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    70 Leu Arg Thr Tyr Asn Leu Arg His Arg Leu Glu Glu Gly Gln Ala Thr
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82 Phe Phe Arg Asp Leu Asp Ile Val Pro Ser Asp Ile Ile Ala Gly Leu

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87		245			250			255
90 Glu Ala	a Asn Asn	Asp Ile	Leu A		Leu Se	er Gly	Met Pro	Val Thr
91	260			265		_	270	
94 Arg Ası	_	Tyr Leu	_	_	Asn Se			Leu Arg
95	275	_		280			285	
	s Glu Val	Cys Tyr		Met Leu	Phe A.		Ala Ala	Tyr Gly
99 29	-		295	T D	- 33- /	300		- Gl- T
102 Trp P:	ro Met Ty	г њеи ме 31		Lys Pr		cys Gly 315	Leu Cy	s Gin Leu 320
105 305 106 Ala A:	ra Ser Cus		-	Len Cv			Pro Ar	
100 AIA A	ig ber cy.	325	3 Cy3	neu cy	330	niu nig	IIO AI	335
110 Pro G	lv Val Th		u Glu	Asp As		Cvs Glv	Cvs As	
111	340			34	_	-11	35	
114 Ala I	le Arg Arg	His Ph	e Leu	Asp Gl	u Asn I	Met Thr	Ala Va	l Asp Ile
115	355	•		360			365	-
118 Val T	yr Thr Se	Cys Hi	s Asp	Ala Va	l Tyr (	Glu Thr	Pro Ph	e Tyr Val
			3,75			380		
122 Ala V	al Asp His	s Asp Ly	s Lys	Lys Va	l Val .	Ile Ser	Ile Ar	g Gly Thr
123 385		39	_			395		400
126 Leu S	er Pro Lys	_	a Leu	Thr As	_	Thr Gly	Asp Al	_
127		405		a 3 m)	410		*** - * -	415
130 Leu P			s His			Leu Gry		
131 134 Val L	420 31 Sor Al:		r Tlo	42	_	Lou Clu	43 (2) (2)	-
134 Val 13	435	d Giu iy	1 116	440	з шуз 1	nea Gra	445	u Mec Vai
138 Leu S		a Phe Gl	v Ara		u Glv A	Ara Glv	_	s His Tvr
	50		455	<b>F</b>		460		2
142 Gly L	eu Ile Va	l Val Gl	y His	Ser Le	u Gly i	Ala Gly	Thr Al	a Ala Ile
143 465		47	0		4	475		480
146 Leu S	er Phe Lei	ı Leu Ar	g Pro	Gln Ty	r Pro !	Thr Leu	Lys Cy	s Phe Ala
147		485			490			495
150 Tyr S		_	y Leu			Asp Ala		
151	500		_ 77_7	50		T 7	51	
154 Lys G	iu Phe va. 515	I THY AL	a vai	vai це 520	u Giy	Lys Asp	ь шец va 525	I PIO AIG
155 158 Tle G		r Gln Le	n Glu		e Ara i	Ara Gln		u Asp Val
	30	C OIN DO	535	Ory III	c mrg .	540		a nop var
		r Thr Lv		Lvs Tr	p Arq	_		y Ala Thr
163 545		55				555		560
	ys Ile Pro	Lys Se	r Glu	Leu Pr	o Glu (	Glu Val	Glu Va	1 Thr Thr
167		565			570			575
170 Leu A	la Ser Th	r Arg Le	u Trp	Thr Hi	s Pro	Ser Asp		r Ile Ala
171	580			58			59	
		r Thr Pr	o Leu		o Pro (	Gly Arg		e His Val
175	595		~-	600			605	
		s Pro Al		GIn Cy	s Cys (			n Glu Glu
179 6	10		615			620	Ĭ	

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	625		_	_		630	_				635	_	_			640
	He	He	Ser	Pro		Met	Leu	His	GIu	His	Leu	Pro	Tyr	Val		Met
187			_	_	645		_	~ 7	_	650	_	_	~3	_	655	
	Glu	Gly	Leu		Lys	Val	Leu	GIu		Tyr	Asn	Lys	GIY	_	Thr	Ala
191				660	_		_		665					670		_
	Leu	Leu		Ala	Ala	Lys	Val		Val	Ser	Pro	Thr		Val	Asp	Leu
195			675		_			680	_				685			
198	Thr		Glu	Leu	Ile	Phe		Gln	Gln	Pro	Leu		Thr	Gly	Pro	Pro
199		690		_			695					700				
		Pro	Thr	Gly	Leu		Leu	Glu	Leu	Pro		Ala	Asp	His	Arg	
	705		_			710		_		_	715					720
	Ser	Ser	Val	Arg		Lys	Ser	Gln	Ser	Glu	Met	Ser	Leu	Glu	_	Phe
207					725				_	730	_	_	_	_	735	
210	Ser	Glu	Gly	_	Leu	Leu	Ser	Pro		Val	Ala	Ala	Ala		Arg	Gln
211				740					745					750		_
214	Asp	Pro	Val	Glu	Leu	Leu	Leu		Ser	Thr	Gln	Glu	_	Leu	Ala	Ala
215			755					760					765			
218	Glu	Leu	G" n	Ma	Arg	Arg		Prc	Leu	Ala	Thr		Glu	Ser	Leu	Sar
219		770					775					780				
222	Asp	Thr	Glu	Ser	Leu	_	Ser	Phe	Asp	Ser	_	Arg	Ser	Ser	Gly	
	785					790					795					800
226	Arg	Ser	Ile	Arg	Gly	Ser	Pro	Ser	Leu	His	Ala	Val	Leu	Glu	Arg	Asp
227					805					810					815	
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	Glu	Val	Gly	Gly		Gly	Gly	Gly	Pro	Ala	Ser	Arg	Gly	Glu		Ala
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251				900	_				905					910		
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259		930		_	_		935				_	940				
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	945	_			_	950					955					960
	Arg	Ala	Gln	Phe		Pro	Asn	Leu	Val	Pro	Lys	Pro	Pro	Arg		Phe
267	_	_		_	965					970			_		975	
	Ala	Gly	Ser		Asp	Pro	Ser	Ser	_	Ile	Ser	Leu	Ser		Ser	Phe
271				980		_			985			_		990		
	Pro	Leu		Ser	Ser	Gly	Glu			t Ası	p Lei	ı Thi			hr G	ly Leu
275			995					100					10			
278 Ser Ser Gln Glu Cys Leu Ala Ala Asp Lys Ile Arg Thr Ser Thr										Fhr						

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378	tcggacactg	agtccctgta	cagcttcgac	tcgcgccgct	cctcaggctt	ccgcagcatc	2520
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			acaggacacg				2700
			ccccagtgct				2760
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			caagccccca				3060
			ctccttcccg				3120
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			ttttggaaaa				4740
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			caaggacacc				5040
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VERIFICATION SUMMARY

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PATENT APPLICATION: US/10/569,224

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